

GenCore version 5.1.6
Copyright (c) 1993 - 2005 CompuGen Ltd.

OM protein - protein search, using sw model

Run on: March 7, 2005, 06:55:26 ; Search time 46.7139 Seconds
(without alignments)
919.008 Million cell updates/sec

Title: US-09-939-537-37

Perfect score: 591

Sequence: 1 TFFSRASBPAYQGGQGNQLY.....LSTPTKDTYDALHMLPPR 111

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 2105692 seqs, 386760381 residues

Total number of hits satisfying chosen parameters: 2105692

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database : A_Geneseq_16Dec04:*

1: geneseqp1980s:*\n2: geneseqp1980s:*\n3: geneseqp2000s:*\n4: geneseqp2001s:*\n5: geneseqp2002s:*\n6: geneseqp2003as:*\n7: geneseqp2003bs:*\n8: geneseqp2004s:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB	ID	Description
1	591	100.0	111	2	AAR78675	Aar78675 T-cell re
2	591	100.0	111	2	AAR89452	Aar89452 Zeta intr
3	583	98.6	142	2	AAR89461	Aar89461 Zeta intr
4	583	98.6	142	2	AAR02226	Aar02226 T-cell re
5	583	98.6	163	2	AAV14196	Aay14196 T-cell re
6	583	98.6	443	6	ABG76488	Abg76488 Humanised
7	583	98.6	443	6	ABG74240	Abg74240 Chimeraic
8	583	98.6	532	2	AAR72728	Aar72728 CD4:gamma
9	583	98.6	532	2	AAR78678	Aar78678 T-cell re
10	583	98.6	532	2	AAR89458	Aar89458 CD4:eta f
11	583	98.6	532	2	AAW02215	AAW02215 CD4:T-cel
12	583	98.6	532	2	AAW63141	AAW63141 Chimeric
13	572	96.8	112	8	ADL67239	Adl67239 Human CD3
14	572	96.8	163	8	ADL67237	Adl67237 Chimeric
15	572	96.8	163	8	ABO84748	AbO84748 Human can
16	572	96.8	357	5	ABG73652	Abg73652 Myvalpha-
17	572	96.8	395	5	ABG73653	Abg73653 Myvalpha-
18	572	96.8	444	2	AAW36845	AAW36845 Single ch
19	572	96.8	473	2	AAW26646	AAW26646 Chimeric
20	572	96.8	514	2	AAW26647	AAW26647 Chimeric
21	572	96.8	532	2	AAW73051	AAW73051 Z33dCH2.z
22	572	96.8	631	7	ADH34696	Adh34696 CEF-spect
23	572	96.8	631	7	AAV84965	Aay84965 Amino ac1
24	572	96.8	634	6	ABR82300	Abb82300 CD19:zeta
25	572	96.8	643	2	AAW73050	Aaw73050 Z33g2G237

26	572	96.8	643	2	AAW73048	Aaw73048 A33 chime
27	572	96.8	651	2	AAW26649	Aaw26649 Chimeric
28	572	96.8	692	2	AAW26650	Aaw26650 Chimeric
29	561.5	95.0	164	8	ABO84747	AbO84747 Human can
30	476.5	80.6	113	3	AAV52340	Aay52340 Full-leng
31	476.5	80.6	113	6	ABU08920	Abu08920 Mouse tru
32	476.5	80.6	113	7	ADH74813	Adh74813 Zeta. 4/2
33	476.5	80.6	113	7	ADH74805	Adh74805 Mouse imm
34	476.5	80.6	164	8	ABO84745	AbO84745 Mouse can
35	476.5	80.6	461	2	AAR85508	Aar85508 Leader-8c
36	476.5	80.6	483	2	AAW82315	Aaw82315 Chimeric
37	472.5	79.9	113	3	AAV52345	Aay52345 Mutant fu
38	472.5	79.9	113	6	ABU08925	Abu08925 Mouse tru
39	472.5	79.9	113	7	ADH74810	Adh74810 CD25/Zeta
40	472.5	79.9	443	2	AAW24027	Aaw24027 Single ch
41	472.5	79.9	443	2	AAW24025	Aaw24025 Single ch
42	467.5	79.1	113	3	AAV52344	Aay52344 Mutant fu
43	467.5	79.1	113	6	ABU08924	Abu08924 Mouse tru
44	467.5	79.1	113	7	ADH74809	Adh74809 CD8/Zeta
45	378.5	64.0	206	2	AAR13273	Aar13273 Murine CD

ALIGNMENTS

RESULT 1	AAW78675	standard; protein; 111 AA.
ID	AAW78675	
XX	AAW78675;	
AC		
XX		
DT	12-APR-1996	(first entry)
XX		
DE	T-cell receptor protein zeta intracellular domain.	
XX		
KW	Chimeric receptor; CD4; T-cell receptor zeta; HIV; cytolysis;	
KM	human immunodeficiency virus; adoptive immunotherapy.	
XX		
OS	Homo sapiens.	
XX		
PN	W09521528-A1.	
XX		
PD	17-AUG-1995.	
XX		
PF	12-JAN-1995;	95MO-US000454.
XX		
PR	14-FEB-1994;	94US-00195395.
XX		
PR	02-AUG-1994;	94US-00284391.
XX		
PA	(GENO) GEN HOSPITAL CORP.	
XX		
XX	Seed B, Banapour B, Romeo C, Kojanus W;	
XX		
DR	WPI; 1995-292893/38.	
XX		
DR	N-PSDB; AAQ96105.	
XX		
PT	Target cytolysis of HIV-infected cells - by chimeric CD4 receptor-bearing cells.	
XX		
PS	Example 10; Fig 27; 118pp; English.	
XX		
CC	The intracellular domain (AAR78675) of human T-cell receptor protein zeta is used in the construction of a chimeric receptor utilized in the	
CC	targeted cytolysis of cells expressing HIV envelope proteins on their	
CC	surface. The chimeric receptor comprises the extracellular domain (pref.	
CC	amino acids 1-394 or 1-200) of CD4 linked to the intracellular portion of	
CC	e.g. zeta	
XX		
SO	Sequence 111 AA;	
XX		
Query Match	100.0%; Score 591; DB 2; Length 111;	
Best Local Similarity	100.0%; Pred. No. 1.6e-59;	
Matches	111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;	

```

QY      1 TRFSRSAPPAYOQOGNOLYNELNGRREYDVLDKRGDPPEMGKPRKKNPOEGLYNE 60
DB      1 TRFSRSAPPAYOQOGNOLYNELNGRREYDVLDKRGDPPEMGKPRKKNPOEGLYNE 60
QY      61 LQDKMAEAYSEIGMKGERRRGKGDGLYQGLSTATKOTYDALHMOALPPR 111
DB      61 LQDKMAEAYSEIGMKGERRRGKGDGLYQGLSTATKOTYDALHMOALPPR 111

RESULT 2
AAR89452 ID AAR89452 standard; peptide; 111 AA.
XX
AC AAR89452;
XX
DT 26-SEP-1996 (first entry)
XX
DE Zeta intracellular domain.
XX
KW CD7; transmembrane domain; chimeric receptor; CD5; CD34; CH2; CH3; IgG1;
KW human; CD4; HIV; proteinaceous alpha-helix; T cell; B cell; neutrophil;
KW dendritic cell; therapy; mammal; infection.
XX
OS Homo sapiens.
XX
PN WO9603883-A1.
XX
PD 15-FEB-1996.
XX
PF 26-JUL-1995; 95WO-US009468.
XX
PR 02-AUG-1994; 94US-00284391.
PR 24-FEB-1995; 95US-00394388.
XX
PA (GENO ) GEN HOSPITAL CORP.
XX
PI Seed B, Banapour B, Romeo C, Kolanus W;
XX
DR WPI; 1996-129034/13.
DR N-PSDB; AAT10799.
XX
PT Membrane-bound chimeric receptor comprising extracellular portion
PT including CD4 fragment - cells expressing receptor can be used for
PT treatment of HIV infection.
XX
PS Example 10; Fig 27; 134pp; English.
XX
CC This sequence represents the zeta intracellular domain. This sequence is
CC included in the membrane bound proteinaceous chimeric receptor of the
CC invention. The extracellular portion of the chimeric receptor contains a
CC fragment of CD4 (amino acids 1-394 or 1-200 of the CD4 sequence see
CC AAR89450 and AAR89451) which specifically recognises and binds HIV-
CC infected cells, but does not mediate HIV infection. The extracellular
CC domain of the receptor is separated from the cell membrane by 48 or 72
CC angstroms, or by one or more proteinaceous alpha-helices. The
CC transmembrane region of the chimeric receptor contains a portion of the
CC CD7 (see AAR89440), CD5 or CD34 transmembrane domain. Alternatively, the
CC extracellular portion of the receptor can also be separated from the
CC intracellular domain by the hinge, CH2 and CH3 domains of human IgG1 (see
CC AAR89441). The cells expressing the receptor are preferably T cells, B
CC cells, neutrophils, or dendritic cells. The therapeutic cells expressing
CC the chimeric receptor are administered to a mammal to treat HIV infection
XX
SQ Sequence 111 AA;
XX
Query Match 100.0%; Score 591; DB 2; Length 111;
Best Local Similarity 100.0%; Pred. No. 1.6e-59;
Matches 111; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY      1 TRFSRSAPPAYOQOGNOLYNELNGRREYDVLDKRGDPPEMGKPRKKNPOEGLYNE 60
DB      1 TRFSRSAPPAYOQOGNOLYNELNGRREYDVLDKRGDPPEMGKPRKKNPOEGLYNE 60

```

```

QY      61 LQDKMAEAYSEIGMKGERRRGKGDGLYQGLSTATKOTYDALHMOALPPR 111
DB      61 LQDKMAEAYSEIGMKGERRRGKGDGLYQGLSTATKOTYDALHMOALPPR 111

RESULT 3
AAR89461 ID AAR89461 standard; peptide; 142 AA.
XX
AC AAR89461;
XX
DT 26-SEP-1996 (first entry)
XX
DE Zeta intracellular domain.
XX
KW CD7; transmembrane domain; chimeric receptor; CD5; CD34; CH2; CH3; IgG1;
KW human; CD4; HIV; proteinaceous alpha-helix; T cell; B cell; neutrophil;
KW dendritic cell; therapy; mammal; infection.
XX
OS Homo sapiens.
XX
PN WO9603883-A1.
XX
PD 15-FEB-1996.
XX
PF 26-JUL-1995; 95WO-US009468.
XX
PR 02-AUG-1994; 94US-00284391.
PR 24-FEB-1995; 95US-00394388.
XX
PA (GENO ) GEN HOSPITAL CORP.
XX
PI Seed B, Banapour B, Romeo C, Kolanus W;
XX
DR WPI; 1996-129034/13.
XX
PT Membrane-bound chimeric receptor comprising extracellular portion
PT including CD4 fragment - cells expressing receptor can be used for
PT treatment of HIV infection.
XX
PS Example 8; Page 96; 134pp; English.
XX
CC This sequence represents the zeta intracellular domain. This sequence is
CC included in the membrane bound proteinaceous chimeric receptor of the
CC invention. The extracellular portion of the chimeric receptor contains a
CC fragment of CD4 (amino acids 1-394 or 1-200 of the CD4 sequence see
CC AAR89450 and AAR89451) which specifically recognises and binds HIV-
CC infected cells, but does not mediate HIV infection. The extracellular
CC domain of the receptor is separated from the cell membrane by 48 or 72
CC angstroms, or by one or more proteinaceous alpha-helices. The
CC transmembrane region of the chimeric receptor contains a portion of the
CC CD7 (see AAR89440), CD5 or CD34 transmembrane domain. Alternatively, the
CC extracellular portion of the receptor can also be separated from the
CC intracellular domain by the hinge, CH2 and CH3 domains of human IgG1 (see
CC AAR89441). The cells expressing the receptor are preferably T cells, B
CC cells, neutrophils, or dendritic cells. The therapeutic cells expressing
CC the chimeric receptor are administered to a mammal to treat HIV infection
XX
SQ Sequence 142 AA;
XX
Query Match 98.6%; Score 583; DB 2; Length 142;
Best Local Similarity 99.1%; Pred. No. 1.8e-58;
Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
QY      2 TRFSRSAPPAYOQOGNOLYNELNGRREYDVLDKRGDPPEMGKPRKKNPOEGLYNE 61
DB      33 TRFSRSAPPAYOQOGNOLYNELNGRREYDVLDKRGDPPEMGKPRKKNPOEGLYNE 92
QY      62 QDKMAEAYSEIGMKGERRRGKGDGLYQGLSTATKOTYDALHMOALPPR 111
DB      93 QDKMAEAYSEIGMKGERRRGKGDGLYQGLSTATKOTYDALHMOALPPR 142

```

Query Match	Best Local Similarity	Score	DB 2	Length	142
Matches 109; Conservative	99.1%;	Pred. No. 1.8e-58;	Mismatches 0;	Indels 0;	Gaps 0
2	FFSRSAEPPAYQOQGNOLYNELNLRREBYDVLDRKRGDPEWGGKPRRRKNPQEGLYNEL	61			
33	KFSRSABPPAYQOQGNOLYNELNLRREBYDVLDRKRGDPEWGGKPRRRKNPQEGLYNEL	92			
62	QKDKAAEYSEIYSGMKGERRRGKGGHGLYGLSTATTQYTDALHMQALLPR	111			
93	QKDKAAEYSEIYSGMKGERRRGKGGHGLYGLSTATTQYTDALHMQALLPR	142			

KW	T cell receptor zeta chain protein; autoimmune disease; diagnosis; SLB;
KM	Systemic lupus erythematosus; therapy.
XX	
OS	Homo sapiens.
XX	
PN	JP11123082-A.
XX	
PD	11-MAY-1999.
XX	
PF	23-OCT-1997; 97JP-00309302.
XX	
PR	23-OCT-1997; 97JP-00309302.
XX	
PA	(TAKE/) TAKEUCHI T.
PA	(NOBE-) NOBEL IGAKU KENKUTSUO YG.
XX	
DR	WPI; 1999-340518/29.
DR	N-PSDB; AAK61131.
XX	
PT	New T cell receptor zeta chain protein - useful for detecting auto-immune
PT	disease.
XX	
PS	Claim 3; Page 12-13; 22pp; Japanese.
XX	
CC	This sequence is the T cell receptor zeta chain protein of the invention,
CC	which is used for detecting an autoimmune disease in a patient. The
CC	protein can be used for detecting an autoimmune disease affliction. Systemic
CC	lupus erythematosus (SLE) can be diagnosed by detecting the mutation of T
CC	cell zeta chain gene and can be treated by complementing the detected
CC	mutation with a normal zeta chain gene or zeta chain protein
XX	
SQ	Sequence 163 AA;
OY	Query Match 98.6%; Score 583; DB 2; Length 163; Best Local Similarity 99.1%; Pred. No. 2,1e-58; Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0
Db	2 RFSRSABPPAYQGCGONQLYNELNGRREYVDLDKRRGRDEMGCKPRKNPOEGLYNEL 61 : : : : : : : : : : : : 54 KFSSRSABPPAYQGCGONQLYNELNGRREYVDLDKRRGRDPMGCKPRKNPOEGLYNEL 113
Oy	62 QKDGMARVYSFIRGKGERRRGKHDLGYOGSLSTATKTQTYDALHMOALPPR 111 : : : : : : : : : : : 114 QKDGMARVYSFIRGKGERRRGKHDLGYOGSLSTATKTQTYDALHMOALPPR 163
Dd	
RESULT 6	
ABG76488	
ID	ABG76488 standard; protein; 443 AA.
XX	
AC	ABG76488;
XX	
DT	13-MAY-2003 (first entry)
XX	
DE	Humanised anti-CEA antibody, hmn14.
XX	
KW	CEA; carcinoembryonic antigen; IGTCR; T-cell receptor; cancer; tumour;
KW	colorectal cancer; breast cancer; lung cancer; hmn14; cytostatic; mouse;
KW	human; zeta signalling chain; CDealpa hinge; humanised antibody.
XX	
OS	Homo sapiens.
OS	Mus sp.
OS	Synthetic.
OS	Chimeric.
XX	
FN	US2002165360-A1.
XX	
PD	07-NOV-2002.
PF	10-DEC-2001; 2001US-00006771.
XX	
PR	30-NOV-2000; 2000US-0250087P.
PR	30-NOV-2000; 2000US-0250090P.

XX (JUNG/) JUNGHANS R P.
XX Junghans RP;
XX WPI; 2003-298705/29.
XX N-PSDB; ABX13168.
XX
XX New chimeric molecule from humanized antibody against carcinoembryonic
XX antigen and having signaling molecules of T cells and other effector
XX cells, useful for the treatment of colorectal, breast and lung cancers.
XX
XX Disclosure; Page 7-8; 20pp; English.
XX
XX The invention relates to a chimeric molecule comprising the
XX carcinoembryonic antigen (CEA) binding domain of humanised antibody MN14
XX as a single chain antibody with a (GGSGS)3 linker, the zeta signalling
XX chain of the T cell receptor (TCR) and an intervening CD8alpha hinge in
XX which the cysteine residues have been mutated, with the IGTcr molecule
XX occupying nucleotides 2426-3766 of the retroviral vector sequence.
XX appearing as ABX13168. The new chimeric molecule expressed in T cells,
XX NK (not defined) or other effector cells are useful in treating patients
XX with cancers expressing the CEA antigen, together with other or with
XX heterologous constructs to engage additional stimulatory and functional
XX properties of the effector cells to enhance the anti-tumour therapeutic
XX efficacy. The cancer disorder includes colorectal, breast and lung
XX cancers. The present sequence represents the chimeric molecule of the
XX invention
XX
XX Sequence 443 AA;
XX
XX
XX Query Match 98.6%; Score 583; DB 6; Length 443;
XX Best Local Similarity 99.1%; Pred. No. 7.3e-58;
XX Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
XX
XX
XX 2 RRSRSAPPAYOQGGONQLYNELNGRREYDVLDRKGRDPEMGKPRKRNPOEGLYNEL 61
XX :|||||
XX 334 KFSRSAPPAYOQGGONQLYNELNGRREYDVLDRKGRDPEMGKPRKRNPOEGLYNEL 393
XX
XX 62 OKDKMAEAYSEIGMKGERRRGKHDELGYGLSTATKDTYDALHMOALPPR 111
XX |||||||
XX 394 OKDKMAEAYSEIGMKGERRRGKHDELGYGLSTATKDTYDALHMOALPPR 443
XX
XX
XX RESULT 7
XX ABG74240
XX ID ABG74240 standard; protein; 443 AA.
XX
XX ABG74240;
XX
XX 23-OCT-2003 (revised)
XX DT 22-APR-2003 (first entry)
XX
XX Chimeric hMN14/T-cell receptor.
XX
XX Retroviral vector; T-cell receptor; hMN14; antibody; IGTcr; receptor;
XX cytostatic; dermatoprotective; neuroprotective; immunostimulant; GD3;
XX ganglioside antigen; MB3.6; PSMA; tumour; 3D8; 4D4; 3B11;
XX prostate-specific membrane antigen; zeta signalling chain; human; cancer;
XX melanoma; neuroendocrine tumour; prostate cancer; small cell lung cancer;
XX mouse; CD8alpha hinge.
XX
XX Homo sapiens.
XX Mus sp.
XX Chimeric.
XX
XX US2002132983-A1.
XX
XX 19-SEP-2002.
XX
XX 10-DEC-2001; 2001US-0006773.
XX
XX 30-NOV-2000; 2000US-0250087P.
XX
XX

PR 30-NOV-2000; 2000US-0250089P.
XX (JUNG/) JUNGHANS R P.
XX Junghans RP;
XX WPI; 2003-208946/20.
XX N-PSDB; ABX16565.
XX
XX New chimeric molecule useful in treating patients with disorders, such as
XX melanoma, neuroendocrine disorders, prostate and small cell lung cancer
XX comprises GD3 and/or PSMA binding domains of antibody.
XX
XX Disclosure; Page 7-8; 35pp; English.
XX
XX The invention relates to a chimeric molecule comprising the GD3
XX (ganglioside antigen) binding domain of antibody MB3.6, with any of 3
XX variable gene sequences, or the PSMA (prostate-specific membrane antigen)
XX binding domain of antibody 3D8, 4D4 and 3B11, with variable gene
XX sequences, the zeta signalling chain of the T cell receptor and an
XX intervening CD8alpha hinge in which cysteine residues have been mutated.
XX The chimeric molecules expressed in T cells or NK cells or other
XX effector cells are useful in treating patients with cancers expressing
XX the GD3 (MB3.6 derivatives) or PSMA antigen (3D8, 4D4, 3B11 derivatives),
XX and/or together with each other or with heterologous constructs to engage
XX additional stimulatory and functional properties of the effector cells to
XX enhance the antitumour therapeutic efficacy (claimed). They are
XX particularly useful in disorders including melanoma, neuroendocrine
XX tumours and prostate and small cell lung cancer. The present sequence is
XX a hMN14 antibody (specific to CEA antigen) in a fusion protein with the
XX modified CD8alpha hinge and the T-cell receptor zeta chain (IGTcr)
XX encoded by a retroviral vector. The hMN14 antibody coding region is
XX replaced with the MB3.6, 3D8, 4D4 or 3B11 genes of the invention.
XX (Updated on 23-Oct-2003 to standardise OS field)
XX
XX Sequence 443 AA;
XX
XX
XX Query Match 98.6%; Score 583; DB 6; Length 443;
XX Best Local Similarity 99.1%; Pred. No. 7.3e-58;
XX Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
XX
XX
XX 2 RRSRSAPPAYOQGGONQLYNELNGRREYDVLDRKGRDPEMGKPRKRNPOEGLYNEL 61
XX :|||||
XX 334 KFSRSAPPAYOQGGONQLYNELNGRREYDVLDRKGRDPEMGKPRKRNPOEGLYNEL 393
XX
XX 62 OKDKMAEAYSEIGMKGERRRGKHDELGYGLSTATKDTYDALHMOALPPR 111
XX |||||||
XX 394 OKDKMAEAYSEIGMKGERRRGKHDELGYGLSTATKDTYDALHMOALPPR 443
XX
XX
XX RESULT 8
XX AAR27278
XX ID AAR27278 standard; protein; 532 AA.
XX
XX AAR27278;
XX
XX 25-MAR-2003 (revised)
XX DT 28-UTL-1995 (first entry)
XX
XX CD4:gamma peptide chimeric protein.
XX
XX Fusion protein; CD4; extracellular domain; zeta; eta; gamma;
XX membrane spanning domain; intracellular domain; type I;
XX integral membrane homodimer; TCR; T cell antigen receptor;
XX extracellular domain; mouse; human; receptor; chimera;
XX HPB-ALL tumour cell line; natural killer cell.
XX
XX Homo sapiens.
XX
XX MO9215322-A1.
XX
XX 17-SEP-1992.
XX
XX

PF 06-MAR-1992; 92WO-US001785.
 XX
 PR 07-MAR-1991; 91US-00665961.
 XX
 PA (GENO) GEN HOSPITAL CORP.
 XX
 PI Seed B, Romeo C, Kolanus W;
 XX
 DR WPI; 1992-331474/40.
 DR N-PSDB; AAQ28706.
 XX
 PT Therapeutic cells expressing chimeric receptors - directing cellular
 PT response to an infective agent, useful in treating HIV-1, AIDS
 PR Pneumocystis carinii infections etc.
 XX
 PS Example 2; Page 74-76; 114pp; English.
 XX
 CC This sequence represents a fusion protein between the CD4 extracellular
 CC domain and the gamma protein membrane spanning domain and intracellular
 CC domain. The Fc-receptor-associated gamma chain is expressed in cell
 CC surface complexes with additional polypeptides, some of which mediate
 CC ligand recognition, and others which have undefined function. Gamma bears
 CC a homodimeric structure and overall organisation very similar to that of
 CC zeta (see also AAQ28704), and is a component of both the mast
 CC cell/basophil high affinity Igb receptor, Fc-epsilon-RI, which consists
 CC of at least three distinct polypeptide chains and one of the low affinity
 CC receptors for IgG, represented in mice by Fc-gamma-RII-alpha. In the
 CC production of the CD4 receptor chimera, the gamma cDNA was isolated from
 CC the HPB-ALL tumour cell line and from human natural killer cells. The
 CC gamma cDNA was joined to the extracellular domain by engineering a BamHI
 CC site just upstream of the membrane spanning domain, by a BamHI site
 CC naturally present a few residues upstream of the membrane spanning
 CC domain. (Updated on 25-MAR-2003 to correct PN field.)
 CC
 SQ Sequence 532 AA;

Query Match 98.6%; Score 583; DB 2; Length 532;
 Best Local Similarity 99.1%; Pred. No. 9, 2e-58;
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 RFSRSAPPPAYOQGNQLYNEINLGRREBYDVLDKRGRDPENWGKPRRRKNPQEGLYNEL 61
 :|||||
 DB 423 KFSRSAPPPAYOQGNQLYNEINLGRREBYDVLDKRGRDPENWGKPRRRKNPQEGLYNEL 482
 QY 62 QCDKMAEAYSEIGMKGERRRGKHGDLVYGLSTATDTYDALHMQALPPR 111
 :|||||
 DB 483 QCDKMAEAYSEIGMKGERRRGKHGDLVYGLSTATDTYDALHMQALPPR 532

RESULT 9

AA878678
 ID AAR78678 standard; protein; 532 AA.

AC AAR78678;
 XX
 DT 16-APR-1996 (first entry)
 XX
 DE T-cell receptor etc.
 XX
 KW Chimeric receptor; CD4; T-cell receptor etc; HIV; cytolysis;
 KW human immunodeficiency virus; adoptive immunotherapy.
 XX
 OS Homo sapiens.
 XX
 PN WO9521528-A1.
 XX
 PD 17-AUG-1995.
 XX
 PF 12-JAN-1995; 95WO-US000454.
 XX
 PR 14-FEB-1994; 94US-00195395.
 PR 02-AUG-1994; 94US-00284391.
 XX

PA (GENO) GEN HOSPITAL CORP.
 XX
 PI Seed B, Banapour B, Romeo C, Kolanus W;
 XX
 DR WPI; 1995-292893/38.
 DR N-PSDB; AAQ96124.
 XX
 PT Target cytolysis of HIV-infected cells - by chimeric CD4 receptor-bearing
 PT cells.
 XX
 PS Example 2; Page 78-79; 118pp; English.
 XX
 CC Fusion proteins comprising the extracellular domain of CD4 fused to T-
 CC cell receptor zeta, gamma or eta (AAR78676-78, respectively) were
 CC expressed in CV1 using a vaccine virus vector. These CD4:zeta, CD4:gamma
 CC and CD4:eta chimeric receptors mediated cytolysis of targets expressing
 CC HIV gp120/41
 XX
 SQ Sequence 532 AA;

Query Match 98.6%; Score 583; DB 2; Length 532;
 Best Local Similarity 99.1%; Pred. No. 9, 2e-58;
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 RFSRSAPPPAYOQGNQLYNEINLGRREBYDVLDKRGRDPENWGKPRRRKNPQEGLYNEL 61
 :|||||
 DB 423 KFSRSAPPPAYOQGNQLYNEINLGRREBYDVLDKRGRDPENWGKPRRRKNPQEGLYNEL 482
 QY 62 QCDKMAEAYSEIGMKGERRRGKHGDLVYGLSTATDTYDALHMQALPPR 111
 :|||||
 DB 483 QCDKMAEAYSEIGMKGERRRGKHGDLVYGLSTATDTYDALHMQALPPR 532

RESULT 10

AA89458
 ID AAR89458 standard; protein; 532 AA.

AC AAR89458;
 XX
 DT 26-SEP-1996 (first entry)
 XX
 DE CD4:eta fusion protein.
 XX
 KW CD7; transmembrane domain; chimeric receptor; CD5; CD34; CH2; CH3; IgG1;
 KW human; CD4; HIV; proteinaceous alpha-helix; T cell; B cell; neutrophil;
 KW dendritic cell; therapy; mammal; infection.
 XX
 OS Synthetic.
 XX
 PN WO9603883-A1.
 XX
 PD 15-FEB-1996.
 XX
 PF 26-JUL-1995; 95WO-US009468.
 XX
 PR 02-AUG-1994; 94US-00284391.
 PR 24-FEB-1995; 95US-00394388.
 XX
 PA (GENO) GEN HOSPITAL CORP.
 XX
 PI Seed B, Banapour B, Romeo C, Kolanus W;
 XX
 DR WPI; 1996-129034/13.
 DR N-PSDB; AAT10803.
 XX
 PT Membrane-bound chimeric receptor comprising extracellular portion
 PT including CD4 fragment - cells expressing receptor can be used for
 PT treatment of HIV infection.
 XX
 PS Example 2; Page 80-81; 134pp; English.
 XX
 CC AAT10801-T10803 represent membrane bound proteinaceous chimeric receptors
 CC of the invention. This sequence represents the CD4:eta chimera. The

CC transmembrane region of the chimeric receptor acts to separate the
 CC intracellular and extracellular domains of the chimera, and contains a
 CC portion of the CD7 (see AAR89440), CD5 or CD34 transmembrane domains.
 CC Alternatively, the extracellular portion of the receptor can be separated
 CC from the intracellular domain by the hinge, CH2 and CH3 domains of human
 CC IgG1 (see AAR89441). The extracellular portion of the chimeric receptor
 CC contains a fragment of CD4 (amino acids 1-394 or 1-200 of the CD4
 CC sequence, see AAR89450 and AAR89451) which specifically recognises and
 CC binds HIV-infected cells, but does not mediate HIV infection. The
 CC extracellular domain of the receptor is separated from the cell membrane
 CC by 48 or 72 angstroms, or by one or more proteinaceous alpha-helices. The
 CC cells expressing the receptor are preferably T cells, B cells,
 CC neutrophils, or dendritic cells. The therapeutic cells expressing the
 CC chimeric receptor are administered to a mammal to treat HIV infection
 XX
 SQ Sequence 532 AA;

Query Match 98.6%; Score 583; DB 2; Length 532;
 Best Local Similarity 99.1%; Pred. No. 9.2e-58;
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 RFSRSABPPAYQOGNOLYNELNGRREYDVLDRGRDPDMGKPRRKPOEGLYNEL 61
 :|||||
 Db 423 KFSRSABPPAYQOGNOLYNELNGRREYDVLDRGRDPDMGKPRRKPOEGLYNEL 482

QY 62 QKDKNAAEAYSEIGMKERRRGKGDGLYOGISTATKDYDLMHQAALPPR 111
 :|||||
 Db 483 QKDKNAAEAYSEIGMKERRRGKGDGLYOGISTATKDYDLMHQAALPPR 532

RESULT 11
 AAM02215
 ID AAM02215 standard; protein; 532 AA.
 AC AAM02215;
 XX
 XX 16-OCT-2003 (revised)
 DT 11-NOV-1996 (first entry)
 XX
 XX CD4:T-cell receptor eta chain chimeric receptor.
 DE
 XX
 XX Chimeric receptor; cellular immunity; adoptive immunotherapy; CD4;
 KW human immunodeficiency virus type 1; HIV-1; AIDS; therapy;
 KW T-cell receptor eta chain; cytotoxic T lymphocyte; CTL.
 XX
 XX Homo; sapiens.
 OS Mus sp.
 OS Chimeric.
 XX
 FH
 FH Key Location/Qualifiers
 FT 1..393
 FT Domain /label= "Extracellular domain"
 FT /note= "CD4 extracellular domain"
 FT 394..396
 FT Region /label= Linker
 FT /note= "encoding DNA contains a BamHI site used for
 FT fusion construction"
 FT 397..532
 FT Region /note= "region of fusion derived from eta chain,
 FT preferred signal-transducing portions for constructs of
 FT the invention are amino acids 421-532, 423-455, 438-455,
 FT 461-494, 494-528 or 400-420"
 FT 400..437
 FT Domain /label= "Transmembrane domain"
 FT /note= "eta chain transmembrane domain"
 FT 438..575
 FT Domain /label= Intracellular domain
 FT /note= "eta chain intracellular domain"
 XX
 XX W09625953-A1.
 PN
 XX
 XX 29-AUG-1996.
 PD
 XX

PF 25-JAN-1996; 96MO-US001056.
 XX
 PR 24-FEB-1995; 95US-00394176.
 XX
 PA (GENO) GEN HOSPITAL CORP.
 XX
 PI Seed B, Romeo C, Kolanus W;
 XX
 XX WPI; 1996-402134/40.
 DR N-PSDB; AAT36760.
 DR
 XX
 XX direction of cellular immune response using therapeutic cell expressing 2
 PT chimeric receptors - comprising region binding to target cell and region
 PT that signals target cell destruction, or CD28 region, partic. for
 PT eliminating HIV-infected cells.
 XX
 XX Claim 7; Page 77-78; 120pp; English.
 PS
 XX A chimeric receptor (AAM0215) comprises the extracellular domain of an
 CC engineered form of the CD4 cellular receptor for HIV and the
 CC transmembrane and intracellular regions, including the cytolitic signal-
 CC transducing portion, of the mouse T-cell receptor eta chain. It can be
 CC obtd. by inserting a gene fusion (AAT36760) into a vaccinia virus vector
 CC and expression in host cells. Chimeric receptors comprising CD4 fused to
 CC eta, eta (see also AAM02213) or Fc receptor gamma (see also AAM02214)
 CC chains are capable of directing cytotoxic T lymphocytes to specifically
 CC recognise and kill cells expressing HIV gp120, thus providing a therapy
 CC for AIDS. (Updated on 16-OCT-2003 to standardise OS field)
 CC
 SQ Sequence 532 AA;

Query Match 98.6%; Score 583; DB 2; Length 532;
 Best Local Similarity 99.1%; Pred. No. 9.2e-58;
 Matches 109; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 RFSRSABPPAYQOGNOLYNELNGRREYDVLDRGRDPDMGKPRRKPOEGLYNEL 61
 :|||||
 Db 423 KFSRSABPPAYQOGNOLYNELNGRREYDVLDRGRDPDMGKPRRKPOEGLYNEL 482

QY 62 QKDKNAAEAYSEIGMKERRRGKGDGLYOGISTATKDYDLMHQAALPPR 111
 :|||||
 Db 483 QKDKNAAEAYSEIGMKERRRGKGDGLYOGISTATKDYDLMHQAALPPR 532

RESULT 12
 AAM83141
 ID AAM83141 standard; protein; 532 AA.
 AC AAM83141;
 XX
 XX 03-FEB-1999 (first entry)
 DT
 XX
 XX Chimeric receptor containing human eta polypeptide.
 DE
 XX
 XX Human; zeta; eta; gamma; membrane-bound chimeric receptor; infection;
 KW tumour; cancer cell; autoimmune-generated cell; T cell receptor; CD3;
 KW CD4; B cell receptor; Fc receptor; pathogen; bacterial; fungal;
 KW protozoan; viral.
 XX
 XX Synthetic.
 OS Homo sapiens.
 OS
 XX US5843728-A.
 PN
 XX 01-DEC-1998.
 PD
 XX 05-APR-1995; 95US-00417495.
 XX
 XX 07-MAR-1991; 91US-00665961.
 PR 06-MAR-1992; 92US-00847566.
 PR 28-FEB-1994; 94US-00203866.
 XX
 XX (GENO) GEN HOSPITAL CORP.
 PA

XX
PI Romeo C, Kolanus W, Seed B;
XX
DR WPI, 1999-044582/04.
XX N-PSDB; AAV70157.
XX
PT Membrane-bound chimeric receptors - comprising extracellular portion
PT which recognises and binds a target cell and an intracellular portion of
XX e.g. a T-cell receptor.
XX
PS Claim 11; Col 45-48; 57pp; English.
XX
CC The present invention describes DNA encoding a membrane-bound chimeric
CC receptor comprising: (a) an extracellular portion that specifically
CC recognises and binds a target cell or a target infective agent; and (b)
CC an intracellular portion of a T-cell receptor CD3, zeta or eta
CC polypeptide, a B-cell receptor polypeptide or an Fc receptor polypeptide.
CC The present sequence represents a chimeric receptor containing the human
CC eta polypeptide. Cells expressing chimeric receptors of the present
CC invention can be administered to mammals in order to destroy pathogens
CC (e.g. bacteria, fungi, protozoa or viruses, especially HIV), cancer cells
CC or autoimmune-generated cells
XX
SQ Sequence 532 AA;

Query Match 98.6%; Score 583; DB 2; Length 532;
Best Local Similarity 99.1%; Pred. No. 9.2e-58; Indels 0; Gaps 0;
Matches 109; Conservative 1; Mismatches 0;

QY 2 RFSRSAPPPAYQGGQNLVNLGRRREYDVLDKRRGRDPENGGKRRKNPQEGLYNEL 61
DB 423 KFSRSAPPPAYQGGQNLVNLGRRREYDVLDKRRGRDPENGGKRRKNPQEGLYNEL 482
QY 62 QKDMMBAVSEIGKERRRGKHDGLYQGLSTATKDTYDALHMOALPPR 111
DB 483 QKDMMBAVSEIGKERRRGKHDGLYQGLSTATKDTYDALHMOALPPR 532

RESULT 13
ADL67239
ID ADL67239 standard; protein; 112 AA.
XX
AC ADL67239;
XX
XX 20-MAY-2004 (first entry)
XX
DE Human CD3 zeta chain intracellular domain.
XX
XX T cell receptor; TCR; CD3 zeta chain; co-stimulatory signalling region;
XX binding element; immunostimulant; therapy; cancer; human.
XX
OS Homo sapiens.
XX
PN US2004043401-A1.
XX
XX 04-MAR-2004.
XX
PD 28-MAY-2003; 2003US-00448256.
XX
PF 28-MAY-2002; 2002US-0383872P.
XX
XX (SLOK) SLOAN KETTERING INST CANCER RES.
XX
PI Sadelain M, Brentjens R, Maher J;
XX
XX WPI, 2004-225696/21.
XX N-PSDB; ADL67228.
XX
XX
PT New nucleic acid polymer encoding a chimeric T cell receptor having a
PT zeta chain portion, useful for treating disorders where the immune
PT response needs to be induced, such as cancer.
XX
XX Disclosure; SEQ ID NO 14; 25pp; English.

XX
CC The invention relates to a nucleic acid polymer encoding a chimeric T
CC cell receptor (TCR) which comprises human CD3 zeta chain intracellular
CC domain, a co-stimulatory signalling region and a binding element that
CC specifically interacts with a selected target. The methods and
CC compositions of the invention are useful for treating disorders where the
CC immune response needs to be induced, such as cancer. The present sequence
CC is human CD3 zeta chain intracellular domain.
XX
SQ Sequence 112 AA;

Query Match 96.8%; Score 572; DB 8; Length 112;
Best Local Similarity 97.3%; Pred. No. 2.4e-57;
Matches 107; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 RFSRSAPPPAYQGGQNLVNLGRRREYDVLDKRRGRDPENGGKRRKNPQEGLYNEL 61
DB 3 KFSRSADAPPAYQGGQNLVNLGRRREYDVLDKRRGRDPENGGKRRKNPQEGLYNEL 62
QY 62 QKDMMBAVSEIGKERRRGKHDGLYQGLSTATKDTYDALHMOALPPR 111
DB 63 QKDMMBAVSEIGKERRRGKHDGLYQGLSTATKDTYDALHMOALPPR 112

RESULT 14
ADL67237
ID ADL67237 standard; protein; 163 AA.
XX
AC ADL67237;
XX
XX 20-MAY-2004 (first entry)
XX
DE Chimeric TCR related human protein #3.
XX
XX T cell receptor; TCR; CD3 zeta chain; co-stimulatory signalling region;
XX binding element; immunostimulant; therapy; cancer; human.
XX
OS Homo sapiens.
XX
PN US2004043401-A1.
XX
XX 04-MAR-2004.
XX
PD 28-MAY-2003; 2003US-00448256.
XX
PF 28-MAY-2002; 2002US-0383872P.
XX
XX (SLOK) SLOAN KETTERING INST CANCER RES.
XX
PI Sadelain M, Brentjens R, Maher J;
XX
XX WPI, 2004-225696/21.
XX
XX
PT New nucleic acid polymer encoding a chimeric T cell receptor having a
PT zeta chain portion, useful for treating disorders where the immune
PT response needs to be induced, such as cancer.
XX
XX Disclosure; SEQ ID NO 12; 25pp; English.
XX
XX
CC The invention relates to a nucleic acid polymer encoding a chimeric T
CC cell receptor (TCR) which comprises human CD3 zeta chain intracellular
CC domain, a co-stimulatory signalling region and a binding element that
CC specifically interacts with a selected target. The methods and
CC compositions of the invention are useful for treating disorders where the
CC immune response needs to be induced, such as cancer. The present sequence
CC is human protein related to the invention.
XX
SQ Sequence 163 AA;

Query Match 96.8%; Score 572; DB 8; Length 163;
Best Local Similarity 97.3%; Pred. No. 3.8e-57;
Matches 107; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

QY 2 RRSRSAPPPAYQOQGNOLYNELNGRREYVDLDRGRDPDMGKPRKKNPOEGLYNEL 61
DB 54 KFSRSADAPAYQOQGNOLYNELNGRREYVDLDRGRDPDMGKPRKKNPOEGLYNEL 113
QY 62 QDKMAEAYSEIGMKERRRGKHDGLYQGLSTATKDTYDALHMQALPPR 111
DB 114 QDKMAEAYSEIGMKERRRGKHDGLYQGLSTATKDTYDALHMQALPPR 163

RESULT 15
AB084748
ID AB084748 standard; protein; 163 AA.
AC AB084748;
XX
XX 18-NOV-2004 (first entry)
XX
XX Human cancer-associated protein HP23-013.2.
XX
XX Human; cancer-associated protein; cytostatic; cancer; leukaemia;
XX Lymphoma; CAP.
XX
XX Homo sapiens.
XX
XX MO2004074320-A2.
XX
XX 02-SEP-2004.
XX
XX 17-FEB-2004; 2004MO-US004730.
XX
XX 14-FEB-2003; 2003US-00367094.
XX 14-MAR-2003; 2003US-00388838.
XX 15-APR-2003; 2003US-00417375.
XX 13-JUN-2003; 2003US-00461862.
XX 15-SEP-2003; 2003US-00663431.
XX 15-DEC-2003; 2003US-00737318.
XX
XX (SAGR-) SAGRES DISCOVERY INC.
XX
XX Morris DW, Morris DW, Malandro MS;
XX
XX WPI; 2004-652914/63.
XX N-PSDB; ABD33068.
XX
XX New isolated cancer-associated polynucleotides and polypeptides useful
XX for diagnosing, preventing or treating cancers, especially lymphoma and
XX leukemia, or in screening for agents that modulate cancer.
XX
XX claim 18; seqid 882; 310pp; English.
XX
XX The invention relates to an isolated nucleic acid comprising at least 10
XX contiguous nucleotides of any of the 233 polynucleotide sequences given
XX in the specification, or its complement. The nucleic acids encode cancer-
XX associated proteins. Also included are an expression vector comprising
XX the isolated nucleic acid cited above, a host cell comprising the above
XX recombinant nucleic acid or expression vector, a microarray for detecting
XX a cancer-associated (CA) nucleic acid comprising at least one probe
XX comprising at least 10 contiguous nucleotides of any of the above-
XX mentioned nucleotide sequences, an isolated polypeptide (encoded within
XX an open reading frame of a CA sequence selected from any of the 95
XX polynucleotide sequences as mentioned in the specification, or its
XX complement), an isolated antibody, (or its antigen binding fragment) that
XX binds to the above polypeptide, a hybridoma that produces the above
XX monoclonal antibody, a pharmaceutical composition comprising the above
XX antibody and a pharmaceutical excipient, a kit for detecting cancer
XX cells (comprising the antibody cited above, methods for diagnosing cancer
XX or for detecting the presence or absence of cancer cells in an
XX individual, a method for inhibiting growth of cancer cells in an
XX individual, a method for delivering a therapeutic agent to cancer cells
XX in an individual, an electronic library comprising the above
XX polynucleotide or polypeptide (or their fragments), methods of screening
XX for anticancer activity or for a bioactive agent capable of modulating
XX the activity of a CA protein (CAP), methods for detecting cancer

```

```

CC associated with expression of a polypeptide in a test cell sample, a
CC method for treating cancers and a method for inhibiting the expression of
CC CA gene in a cell. The composition and methods are useful for detecting,
CC diagnosing, preventing and treating cancers, especially lymphoma and
CC leukaemia. These may also be used in screening for agents that modulate
CC cancer. The present sequence is a human CAP protein sequence. Note: The
CC sequence data for this patent did not form part of the printed
CC specification, but was obtained in electronic format directly from WIPO
CC at ftp.wipo.int/pub/published_pct_sequences
XX
XX Sequence 163 AA;

```

```

Query Match          96.8%; Score 572; DB 8; Length 163;
Best Local Similarity 97.3%; Pred. No. 3.8e-57;
Matches 107; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY 2 RRSRSAPPPAYQOQGNOLYNELNGRREYVDLDRGRDPDMGKPRKKNPOEGLYNEL 61
DB 54 KFSRSADAPAYQOQGNOLYNELNGRREYVDLDRGRDPDMGKPRKKNPOEGLYNEL 113
QY 62 QDKMAEAYSEIGMKERRRGKHDGLYQGLSTATKDTYDALHMQALPPR 111
DB 114 QDKMAEAYSEIGMKERRRGKHDGLYQGLSTATKDTYDALHMQALPPR 163

```

```

Search completed: March 7, 2005, 07:13:07
Job time : 48.7139 secs

```